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INDEX 'ADISALERTS, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, AQUASCI, BIOBUSINESS, BIOCOMMERCE, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CANCERLIT, CAPLUS, CEABA-VTB, CEN, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DRUGB, DRUGLAUNCH, DRUGMONOG2, ...' ENTERED AT 09:52:52 ON 14 NOV 2001

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NEWS 6 Apr 23 PRE-1967 REFERENCES NOW SEARCHABLE IN CAPLUS AND CA
NEWS 7 May 07 DGENE Reload
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0.15

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39 FILES HAVE ONE OR MORE ANSWERS, 62 FILES SEARCHED IN STNINDEX

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F39	1	NAPRALERT

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L2 13541 EUKARYO? AND INITI? AND FACTO? AND KINAS?

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17 FILES SEARCHED...

L3 7289 EIF? AND KINAS?

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18 FILES SEARCHED...
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PROCESSING IS APPROXIMATELY 84% COMPLETE FOR L3
PROCESSING IS APPROXIMATELY 98% COMPLETE FOR L3
PROCESSING COMPLETED FOR L3
L5 2528 DUP REM L3 (4761 DUPLICATES REMOVED)

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PROCESSING IS APPROXIMATELY 18% COMPLETE FOR L2
PROCESSING IS APPROXIMATELY 24% COMPLETE FOR L2
PROCESSING IS APPROXIMATELY 30% COMPLETE FOR L2
PROCESSING IS APPROXIMATELY 36% COMPLETE FOR L2
PROCESSING IS APPROXIMATELY 42% COMPLETE FOR L2
PROCESSING IS APPROXIMATELY 48% COMPLETE FOR L2
PROCESSING IS APPROXIMATELY 54% COMPLETE FOR L2
PROCESSING IS APPROXIMATELY 67% COMPLETE FOR L2
PROCESSING IS APPROXIMATELY 77% COMPLETE FOR L2
PROCESSING IS APPROXIMATELY 85% COMPLETE FOR L2
PROCESSING IS APPROXIMATELY 96% COMPLETE FOR L2
PROCESSING COMPLETED FOR L2
L6 10028 DUP REM L2 (3513 DUPLICATES REMOVED)

=> d hi
'HI' IS NOT A VALID FORMAT
In a multifile environment, a format can only be used if it is valid
in at least one of the files. Refer to file specific help messages
or the STNGUIDE file for information on formats available in
individual files.
REENTER DISPLAY FORMAT FOR ALL FILES (FILEDEFAULT):end

=> d his

(FILE 'HOME' ENTERED AT 09:52:28 ON 14 NOV 2001)

INDEX 'ADISALERTS, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, AQUASCI, BIOBUSINESS, BIOCOMMERCE, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CANCERLIT, CAPLUS, CEABA-VTB, CEN, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DRUGB, DRUGLAUNCH, DRUGMONOG2, ...' ENTERED AT 09:52:52 ON 14 NOV 2001

SEA (EUKARYO? AND INITI? AND FACTO? AND KINAS?) OR (EIF-2? AND

1 FILE ADISALERTS
39 FILE AGRICOLA
5 FILE AQUASCI
2 FILE BIOBUSINESS
1 FILE BIOCOMMERCE
912 FILE BIOSIS
9 FILE BIOTECHABS
9 FILE BIOTECHDS
501 FILE BIOTECHNO
25 FILE CABA
548 FILE CANCERLIT
0* FILE CAPLUS
1 FILE CEABA-VTB
2 FILE CEN
23 FILE CONFSCI
14 FILE DDFU
160 FILE DGENE
1 FILE DRUGNL
21 FILE DRUGU
14 FILE EMBAL
673 FILE EMBASE
795 FILE ESBIODBASE
1 FILE FROSTI
1 FILE FSTA
0* FILE GENBANK
0* FILE IFIPAT
11 FILE JICST-EPLUS
463 FILE LIFESCI
1256 FILE MEDLINE
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0* FILE NTIS
0* FILE PASCAL
1 FILE PHIN
0* FILE PROMT
923 FILE SCISEARCH
0* FILE TOXLIT
0* FILE USPATFULL
0* FILE WPIDS
0* FILE WPINDEX
1 FILE NAPRALERT
0* FILE NLDB

SEA EUKARYO? AND INITI? AND FACTO? AND KINAS?

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9 FILE BIOTECHDS
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659 FILE CAPLUS
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 6 FILE CONFSCI
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 520 FILE EMBASE
 746 FILE ESBIODBASE
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 1 FILE FSTA
 491 FILE GENBANK
 10 FILE IFIPAT
 10 FILE JICST-EPLUS
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 620 FILE MEDLINE
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 116 FILE PASCAL
 1 FILE PHIN
 7 FILE PROMT
 693 FILE SCISEARCH
 116 FILE TOXLIT
 7669 FILE USPATFULL
 11 FILE WPIDS
 11 FILE WPINDEX
 1 FILE NAPRALERT
 19 FILE NLDB

L1 QUE EUKARYO? AND INITI? AND FACTO? AND KINAS?

FILE 'USPATFULL, ESBIODBASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE,
 GENBANK, BIOTECHNO, LIFESCI, CANCERLIT, DGENE, PASCAL, TOXLIT, AGRICOLA,
 CABA, NLDB, DRUGU, EMBAL, WPIDS, IFIPAT, JICST-EPLUS' ENTERED AT 10:48:09
 ON 14 NOV 2001

L2 13541 S EUKARYO? AND INITI? AND FACTO? AND KINAS?
 L3 7289 S EIF? AND KINAS?
 L4 9902 S L2 NOT L3
 L5 2528 DUP REM L3 (4761 DUPLICATES REMOVED)
 L6 10028 DUP REM L2 (3513 DUPLICATES REMOVED)

=> s 15 and 14790?
 5 FILES SEARCHED...
 17 FILES SEARCHED...
 L7 1 L5 AND 14790?

=> d ti

L7 ANSWER 1 OF 1 GENBANK.RTM. COPYRIGHT 2001

TITLE (TI): Direct Submission

=> d 17

L7 ANSWER 1 OF 1 GENBANK.RTM. COPYRIGHT 2001

LOCUS (LOC): SCCHRXVI GenBank (R)
 GenBank ACC. NO. (GBN): Z71255
 CAS REGISTRY NO. (RN): 175825-73-7
 SEQUENCE LENGTH (SQL): 165536
 MOLECULE TYPE (CI): DNA; linear
 DIVISION CODE (CI): Plants, fungi, algae

DATE (DATE): 11 Oct 2000
 DEFINITION (DEF): S.cerevisiae chromosome XVI 165536 bp sequence, cen
 rightwards.
 KEYWORDS (ST): ARO7; ATPase; beta-transducin; CCL1; CDC21 homologue;
 CDE1; CDE2; CDE3; centromere; CHL1; chorismate mutase;
 citrate synthase; coatomer zeta subunit; cyclin; delta
 element; DnaJ; DSS4; electro; transport flavoprotein;
 eukaryotic initiation factor 5; FlFo-AT; synthase g
 subunit; gamma adaptin; GLN1; glutamine synthetase;
 Grave's disease; guanine-releasing protein; HAL1;
 halotolerance; histidine tRNA synthetase; HTS1;
 isocitrate lyase; LPT1; LTR; MAK3; MFS1; Mitochondrial
 carrier protein; N-acetyltransferase; NADPH-cytochrome
 P450 reductase; negative regulator; NHP6A; nonhistone
 chromosomal protein 6A; OSD1; phenylalanyl-tRN;
 synthetase; phosphotyrosine protein phosphatase;
 polyposis; protamine P1; protein **kinase**;
 protein transport; regulator; protein; ribosomal
 protein L37a; RNA polymerase I; ROX1; RPA2; SEC8; SMK1;
 spermidine synthase; TIF5; TKL1; transfer RNA-Gly;
 transfer RNA-Lys; transfer RNA-Phe; transfer RNA-Ser;
 transketolase; tropomyosin; ubiquitin-activating
 enzyme; vacuola; H(+)-ATPase 54 kda subunit; VMA13;
 YMC1; YME1; YTA11; zinc finger; protein
 SOURCE: baker's yeast.
 ORGANISM (ORGN): Saccharomyces cerevisiae
 Eukaryota; Fungi; Ascomycota; Saccharomycotina;
 Saccharomycetes; Saccharomycetales; Saccharomycetaceae;
 Saccharomyces

NUCLEIC ACID COUNT (NA): 52201 a 31306 c 31323 g 50706 t

COMMENT:

SC8132X Z48483 1 to 25540
 SC9723 Z48951 100 to 17660
 SC9531X Z49919 104 to 27166
 SC9367 Z49274 10 to 42190
 SC3085 Z68111 790 to 10339 (PCR product)
 SC9499X Z49219 134 to 43776

The most significant matches using FASTA -o to a non-redundant
 database compiled from Swissprot 32 and PIR 46 and Wormpep9 are
 included.

All CDS over 100 codons have been analysed. CDS that are largely or
 completely overlapped by a larger CDS have been omitted from this
 analysis.

The more significant matches with motifs in the PROSITE database
 are also included but some of these may be fortuitous. The length
 in codons/amino-acids is given for each CDS as is the calculated
 codon adaptation index (CAI).

tRNA genes and LTRs are also included.

Notes:

This sequence has been compiled from the following submissions of
 individual contigs.

REFERENCE: 1 (bases 1 to 165536)
 AUTHOR (AU): Badcock, K.; Bowman, S.; Churcher, C.M.; Pearson, D.;
 Rajandream, M.A.; Walsh, S.V.; Barrell, B.G.
 TITLE (TI): Direct Submission
 JOURNAL (SO): Submitted (19-APR-1996) Saccharomyces cerevisiae
 chromosome IV sequencing project, Sanger Centre,
 Hinxton Hall, Hinxton, Cambridge CB10 1RQ. E-mail:
 barrell@sanger.ac.uk

FEATURES (FEAT):

Feature Key	Location	Qualifier
=====	+	=====

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 KPTRKGRNSSYGVLNRVSRVETRWDAVEDRFND
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 42)
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 and PS00339 Aminoacyl-transfer RNA
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 APKKGKLQVSLKTPKGT KDWADSDMVIREAIFST
 LSGLFKKHGGVTIDTPVFELREIL
 AGKYGEDSKLIYNLEDQG GELCSLRYDLTVPFAR

<-----User Break----->

u
 => d his

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 DDFU, DGENE, DRUGB, DRUGLAUNCH, DRUGMONOG2, ...' ENTERED AT 09:52:52 ON
 14 NOV 2001

SEA (EUKARYO? AND INITI? AND FACTO? AND KINAS?) OR (EIF-2? AND

 1 FILE ADISALERTS
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 912 FILE BIOSIS
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 14 FILE EMBAL
 673 FILE EMBASE
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FILE 'USPATFULL, ESBIODASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE,
GENBANK, BIOTECHNO, LIFESCI, CANCERLIT, DGENE, PASCAL, TOXLIT, AGRICOLA,
CABA, NLDB, DRUGU, EMBAL, WPIDS, IFIPAT, JICST-EPLUS' ENTERED AT 10:48:09
ON 14 NOV 2001

L2 13541 S EUKARYO? AND INITI? AND FACTO? AND KINAS?
L3 7289 S EIF? AND KINAS?
L4 9902 S L2 NOT L3
L5 2528 DUP REM L3 (4761 DUPLICATES REMOVED)
L6 10028 DUP REM L2 (3513 DUPLICATES REMOVED)
L7 1 S L5 AND 14790?

=> s l5 and human? and recombina?

3 FILES SEARCHED...

7 FILES SEARCHED...

11 FILES SEARCHED...

13 FILES SEARCHED...

L8 240 L5 AND HUMAN? AND RECOMBIN?

=> s l6 and human? and recombina?

COMMAND INTERRUPTED

REENTER FILE 'USPATFULL, ESBIODASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE,
GENBANK, BIOTECHNO, LIFESCI, CANCERLIT, DGENE, PASCAL, TOXLIT, AGRICOLA, CABA,
NLDB, DRUGU, EMBAL, WPIDS, IFIPAT, JICST-EPLUS'
AND TRY AGAIN, OR ENTER '?' FOR MORE INFORMATION.

QUERY PROCESSING INTERRUPTED

The system was temporarily unable to process your query. If this
message appears repeatedly, please notify the Help Desk. Enter "HELP
STN" for information on contacting the nearest STN Help Desk by
telephone or via SEND in the STNMAIL file.

=> s l6 and human? and recombina?

THIS COMMAND NOT AVAILABLE IN THE CURRENT FILE

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command can only be used to look at the index in a file which has an
index. Enter "HELP COMMANDS" at an arrow prompt (=>) for a list of
commands which can be used in this file.

=> FIL F1-F24

COST IN U.S. DOLLARS

SINCE FILE	TOTAL
ENTRY	SESSION
125.75	167.30

FULL ESTIMATED COST

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Enter "HELP STN" for information on contacting the nearest STN Help
Desk by telephone or via SEND in the STNMAIL file.

=> d his

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SEA (EUKARYO? AND INITI? AND FACTO? AND KINAS?) OR (EIF-2? AND

1	FILE ADISALERTS
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520  FILE EMBASE
746  FILE ESBIODBASE
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491  FILE GENBANK
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334  FILE LIFESCI
620  FILE MEDLINE
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116  FILE PASCAL
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    7  FILE PROMT
693  FILE SCISEARCH
116  FILE TOXLIT
7669 FILE USPATFULL
    11 FILE WPIDS
    11 FILE WPINDEX
    1  FILE NAPRALERT
    19 FILE NLDB

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L1 QUE EUKARYO? AND INITI? AND FACTO? AND KINAS?

FILE 'USPATFULL, ESBIODBASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE,
GENBANK, BIOTECHNO, LIFESCI, CANCERLIT, DGENE, PASCAL, TOXLIT, AGRICOLA,
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L2      13541 S EUKARYO? AND INITI? AND FACTO? AND KINAS?
L3      7289 S EIF? AND KINAS?
L4      9902 S L2 NOT L3
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L6      10028 DUP REM L2 (3513 DUPLICATES REMOVED)
L7      1 S L5 AND 14790?
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FILE 'USPATFULL, ESBIODBASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE,
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=> s l6 and human? and recombina?

COMMAND INTERRUPTED

REENTER FILE 'USPATFULL, ESBIODBASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE,
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QUERY PROCESSING INTERRUPTED

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=> s l8 and alpha?

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=> FIL F1-F24

COST IN U.S. DOLLARS

FULL ESTIMATED COST

SINCE FILE	TOTAL
ENTRY	SESSION
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- L8 ANSWER 198 OF 240 MEDLINE
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- L8 ANSWER 212 OF 240 MEDLINE
TI Enhancement by retinoic acid and dibutyryl cyclic adenosine 3':5'-monophosphate of the differentiation and gene expression of **human** neuroblastoma cells induced by interferon.
- L8 ANSWER 213 OF 240 MEDLINE
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- L8 ANSWER 214 OF 240 MEDLINE
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- L8 ANSWER 215 OF 240 GENBANK.RTM. COPYRIGHT 2001
TITLE (TI): Complete genome sequence of an aerobic hyper-thermophilic crenarchaeon, Aeropyrum pernix K1
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- L8 ANSWER 216 OF 240 GENBANK.RTM. COPYRIGHT 2001

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TI TNF-binding protein ameliorates inhibition of skeletal muscle protein
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TI THE INTERFERON-INDUCED PROTEIN P1/**EIF-2** ALPHA **KINASE**.

L8 ANSWER 220 OF 240 DGENE COPYRIGHT 2001 DERWENT INFORMATION LTD

TI Enhancing expression of nucleic acids in cells - by using modified
vectors which comprise the nucleic acid and also nucleic acid encoding a
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TI Regulation of gene expression by mRNA splicing is carried out using a
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L8 ANSWER 234 OF 240 DGENE COPYRIGHT 2001 DERWENT INFORMATION LTD
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TI EUROPEAN PATENT DISCLOSURES PRIVATE.

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- L9 ANSWER 141 OF 177 CAPLUS COPYRIGHT 2001 ACS
 TI Molecular mechanisms of interferon action. Phosphorylation of protein P1 and protein synthesis initiation factor **eIF-2.alpha**. in untreated and interferon-treated cells
- L9 ANSWER 142 OF 177 MEDLINE
 TI FANCC interacts with Hsp70 to protect hematopoietic cells from IFN-gamma/TNF-**alpha**-mediated cytotoxicity.
- L9 ANSWER 143 OF 177 MEDLINE
 TI Latently expressed **human** herpesvirus 8-encoded interferon regulatory factor 2 inhibits double-stranded RNA-activated protein **kinase**.
- L9 ANSWER 144 OF 177 MEDLINE
 TI Mouse interferon-inducible RNA-dependent protein **kinase** Pkr gene: cloning and sequence of the 5'-flanking region and functional

identification of the minimal inducible promoter.

- L9 ANSWER 145 OF 177 MEDLINE
TI Activation of NF-kappa B by the dsRNA-dependent protein **kinase**, PKR involves the I kappa B **kinase** complex.
- L9 ANSWER 146 OF 177 MEDLINE
TI Analysis of transcription factors regulating induction of indoleamine 2,3-dioxygenase by IFN-gamma.
- L9 ANSWER 147 OF 177 MEDLINE
TI Mutations in the protein **kinase**-binding domain of the NS5A protein in patients infected with hepatitis C virus type 1a are associated with treatment response.
- L9 ANSWER 148 OF 177 MEDLINE
TI JNK2 and IKKbeta are required for activating the innate response to viral infection.
- L9 ANSWER 149 OF 177 MEDLINE
TI **Human** immunodeficiency virus type 1 tat protein activates transcription factor NF-kappaB through the cellular interferon-inducible, double-stranded RNA-dependent protein **kinase**, PKR.
- L9 ANSWER 150 OF 177 MEDLINE
TI Repression of the PKR protein **kinase** by the hepatitis C virus NS5A protein: a potential mechanism of interferon resistance.
- L9 ANSWER 151 OF 177 MEDLINE
TI The orf virus OV20.0L gene product is involved in interferon resistance and inhibits an interferon-inducible, double-stranded RNA-dependent **kinase**.
- L9 ANSWER 152 OF 177 MEDLINE
TI Interferon regulatory factor 3 and CREB-binding protein/p300 are subunits of double-stranded RNA-activated transcription factor DRAF1.
- L9 ANSWER 153 OF 177 MEDLINE
TI Double-stranded RNA-dependent protein **kinase** mediates c-Myc suppression induced by type I interferons.
- L9 ANSWER 154 OF 177 MEDLINE
TI Nuclear localization of the interferon-inducible protein **kinase** PKR in **human** cells and transfected mouse cells.
- L9 ANSWER 155 OF 177 MEDLINE
TI Activation of the interferon-inducible enzymes, 2',5'-oligoadenylate synthetase and PKR by **human** T-cell leukemia virus type I Rex-response element.
- L9 ANSWER 156 OF 177 MEDLINE
TI Role of the amino-terminal residues of the interferon-induced protein **kinase** in its activation by double-stranded RNA and heparin.
- L9 ANSWER 157 OF 177 MEDLINE
TI Double-stranded RNA-dependent protein **kinase** activates transcription factor NF-kappa B by phosphorylating I kappa B.
- L9 ANSWER 158 OF 177 MEDLINE
TI Hepatitis delta virus replication in vitro is not affected by interferon-**alpha** or -gamma despite intact cellular responses to interferon and dsRNA.

L9 ANSWER 159 OF 177 MEDLINE
 TI Enhancement by retinoic acid and dibutyryl cyclic adenosine 3':5'-monophosphate of the differentiation and gene expression of **human** neuroblastoma cells induced by interferon.

L9 ANSWER 160 OF 177 MEDLINE
 TI Mechanism of interferon action. Interferon **alpha** inhibits vesicular stomatitis virus primary transcript accumulation in P1/**eIF-2 alpha** protein **kinase**-deficient **human** fibroblast cells.

L9 ANSWER 161 OF 177 MEDLINE
 TI Functional properties of proteins coded by three **human alpha**-interferon genes and a pseudogene.

L9 ANSWER 162 OF 177 GENBANK.RTM. COPYRIGHT 2001
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L9 ANSWER 163 OF 177 GENBANK.RTM. COPYRIGHT 2001
 TITLE (TI): Complete Sequence and Gene Organization of the Genome of a Hyper-thermophilic Archaeobacterium, *Pyrococcus horikoshii* OT3
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L9 ANSWER 164 OF 177 CANCERLIT
 TI THE INTERFERON-INDUCED PROTEIN P1/**EIF-2 ALPHA KINASE**.

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 TI Regulation of gene expression by mRNA splicing is carried out using a cis-acting nucleotide sequence controlled by phosphorylation of the **alpha**-subunit of eukaryotic initiation factor 2 -

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 TI Regulation of gene expression by mRNA splicing is carried out using a cis-acting nucleotide sequence controlled by phosphorylation of the **alpha**-subunit of eukaryotic initiation factor 2 -

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 TI Regulation of gene expression by mRNA splicing is carried out using a cis-acting nucleotide sequence controlled by phosphorylation of the **alpha**-subunit of eukaryotic initiation factor 2 -

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 TI Regulation of gene expression by mRNA splicing is carried out using a cis-acting nucleotide sequence controlled by phosphorylation of the **alpha**-subunit of eukaryotic initiation factor 2 -

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 TI Regulation of gene expression by mRNA splicing is carried out using a cis-acting nucleotide sequence controlled by phosphorylation of the **alpha**-subunit of eukaryotic initiation factor 2 -

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 TI Regulation of gene expression by mRNA splicing is carried out using a cis-acting nucleotide sequence controlled by phosphorylation of the **alpha**-subunit of eukaryotic initiation factor 2 -

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TI Regulation of gene expression by mRNA splicing is carried out using a
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TI Regulation of gene expression by mRNA splicing is carried out using a
cis-acting nucleotide sequence controlled by phosphorylation of the
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TI Regulation of gene expression by mRNA splicing is carried out using a
cis-acting nucleotide sequence controlled by phosphorylation of the
alpha-subunit of eukaryotic initiation factor 2 -

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TI Parasitology (Plasmodium falciparum) Protein **Kinase** Gene May Be
Involved in Erythrocyte Rupture

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TI Malaria (Biochemistry) "Molecular Cloning, Characterization and
Localization of PfPK4, an **eIF-2 alpha Kinase**
-Related Enzyme from the Malarial Parasite Plasmodium falciparum."

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TI Interferons: biochemistry and mechanisms of action.

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TI Interferons and their Action.

=> d his

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DDFU, DGENE, DRUGB, DRUGLAUNCH, DRUGMONOG2, ...' ENTERED AT 09:52:52 ON
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L2 13541 S EUKARYO? AND INITI? AND FACTO? AND KINAS?
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L8 240 S L5 AND HUMAN? AND RECOMBIN?

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ON 14 NOV 2001

L9 177 S L8 AND ALPH?

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COMMAND INTERRUPTED

REENTER FILE 'USPATFULL, ESBIODBASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE,
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AND TRY AGAIN, OR ENTER '?' FOR MORE INFORMATION.

QUERY PROCESSING INTERRUPTED

The system was temporarily unable to process your query. If this
message appears repeatedly, please notify the Help Desk. Enter "HELP
STN" for information on contacting the nearest STN Help Desk by
telephone or via SEND in the STNMAIL file.

=> help

The arrow (=>) is the system prompt, where you enter a command. For
an explanation of system commands, files, formats, etc., enter "HELP"
and the name of the item you want explained at an arrow prompt (=>).
Enter "HELP COMMANDS" for a list of commands that can be used in this
file. Enter "HELP MESSAGES" for a list of online explanations that
are available. The "?" can be used as a synonym for "HELP".

Help is also available at any prompt, and after any error message.
Enter "HELP" or "?" at a prompt to see an explanation of the options.
After an error message, enter "HELP" or "?" at the next prompt and you
will receive a more detailed explanation of the error and how to
correct it.

Automatic help is also available. When AUHELP is 'ON', you will
automatically receive help following an error message. For more
information on AUHELP, enter "HELP SET AUHELP" at an arrow
prompt (=>).

Users who need additional assistance can contact the Help Desk at
their nearest STN Service Center. Enter "HELP STN" for information on
STN Service Centers. You may also choose to contact the database
representative for the file you are searching, for more detailed help
on database content and search strategy. For information on how to
contact database representatives for the current file, enter "HELP
DESK" at an arrow prompt (=>).